

FIGURE 1A

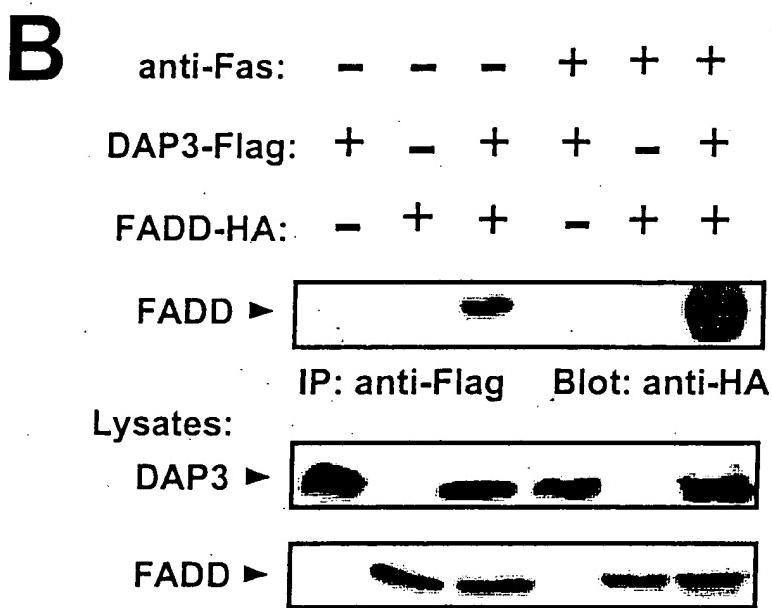


FIGURE 1B

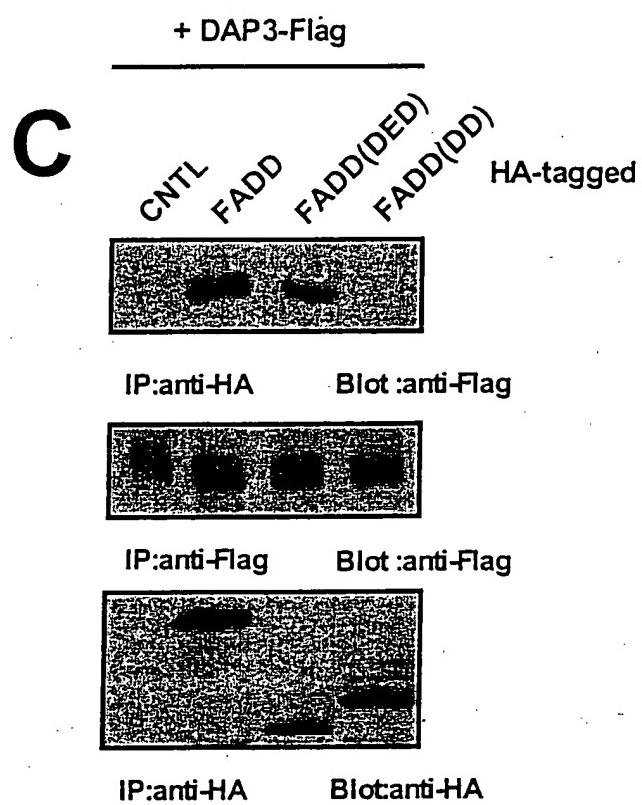


FIGURE 1C

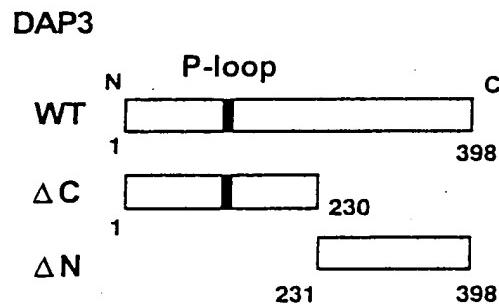
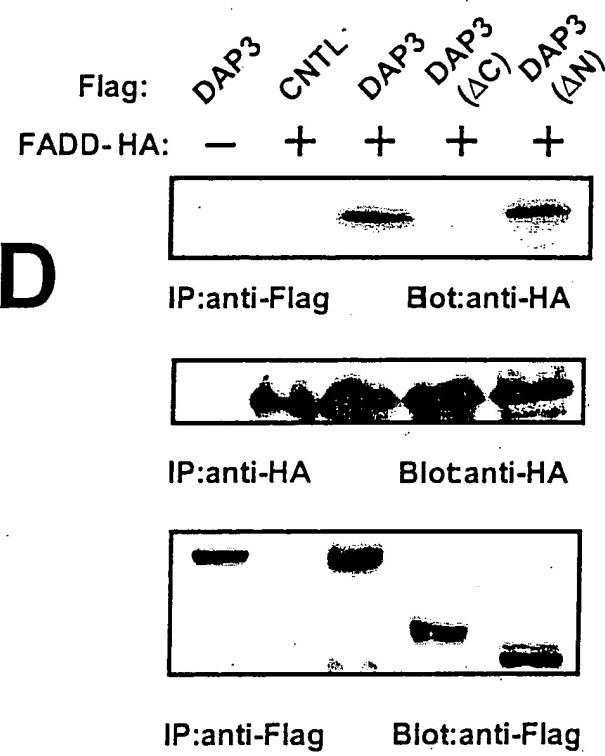


FIGURE 1D

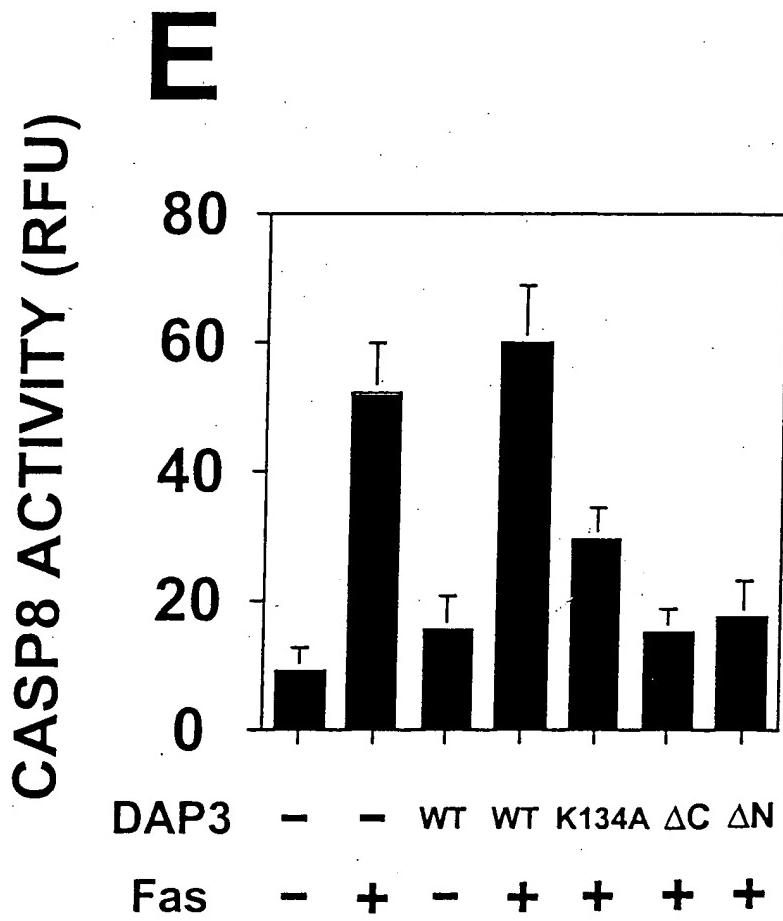


FIGURE 1E

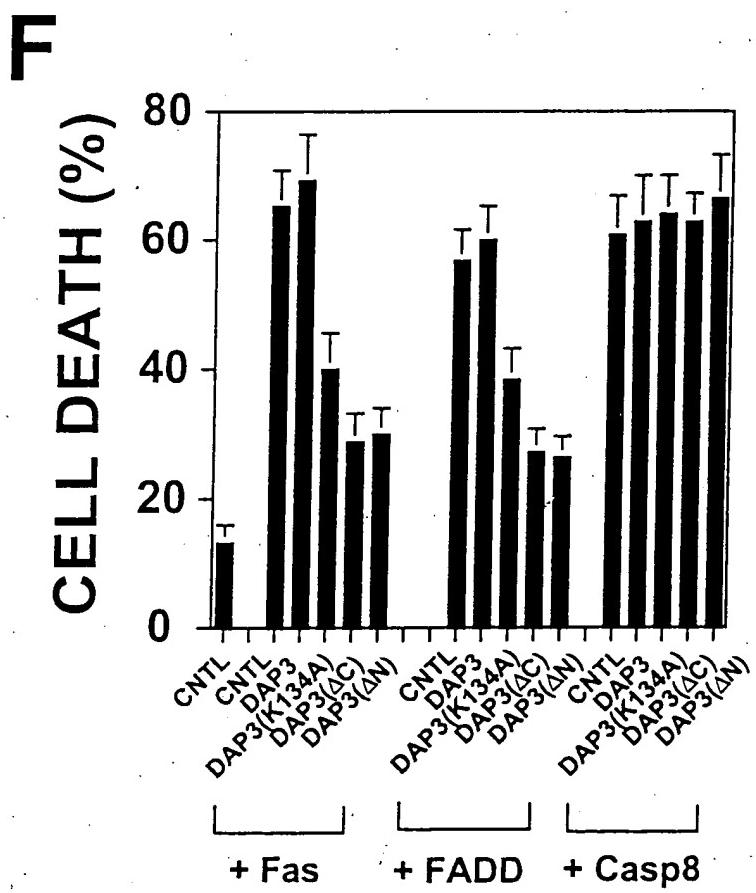


FIGURE 1F

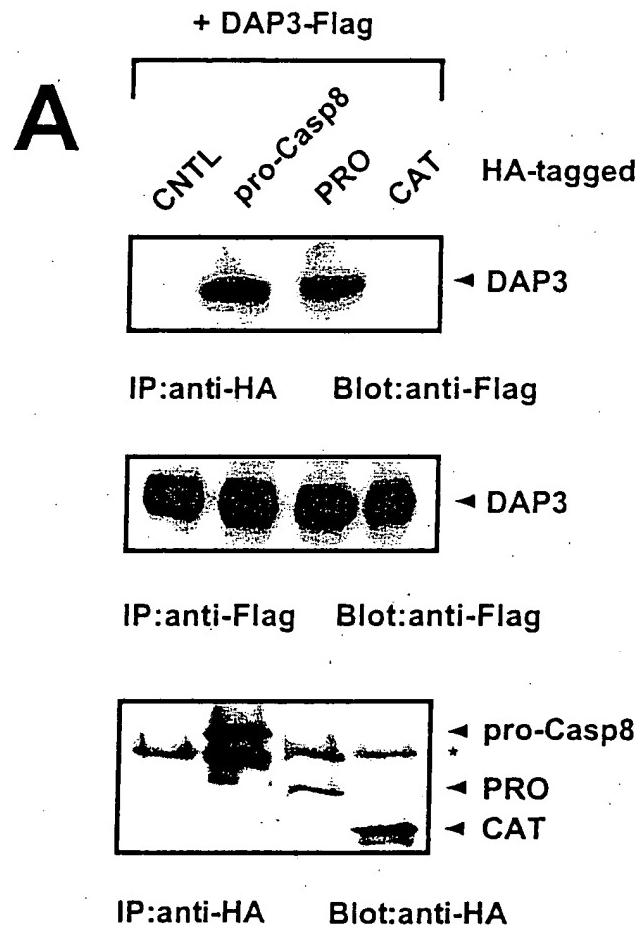


FIGURE 2A

B

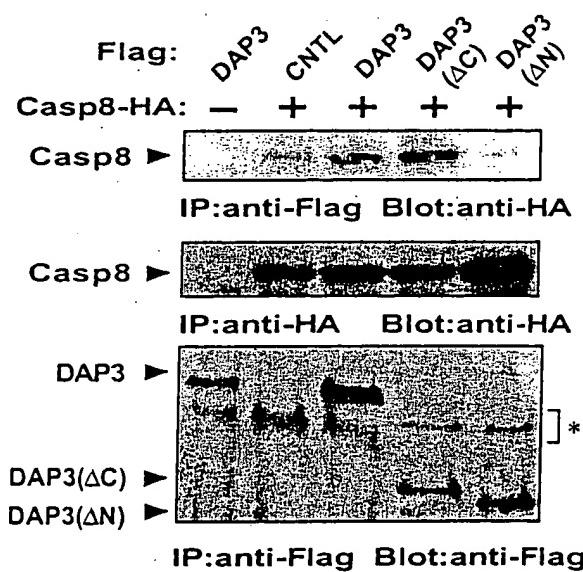


FIGURE 2B

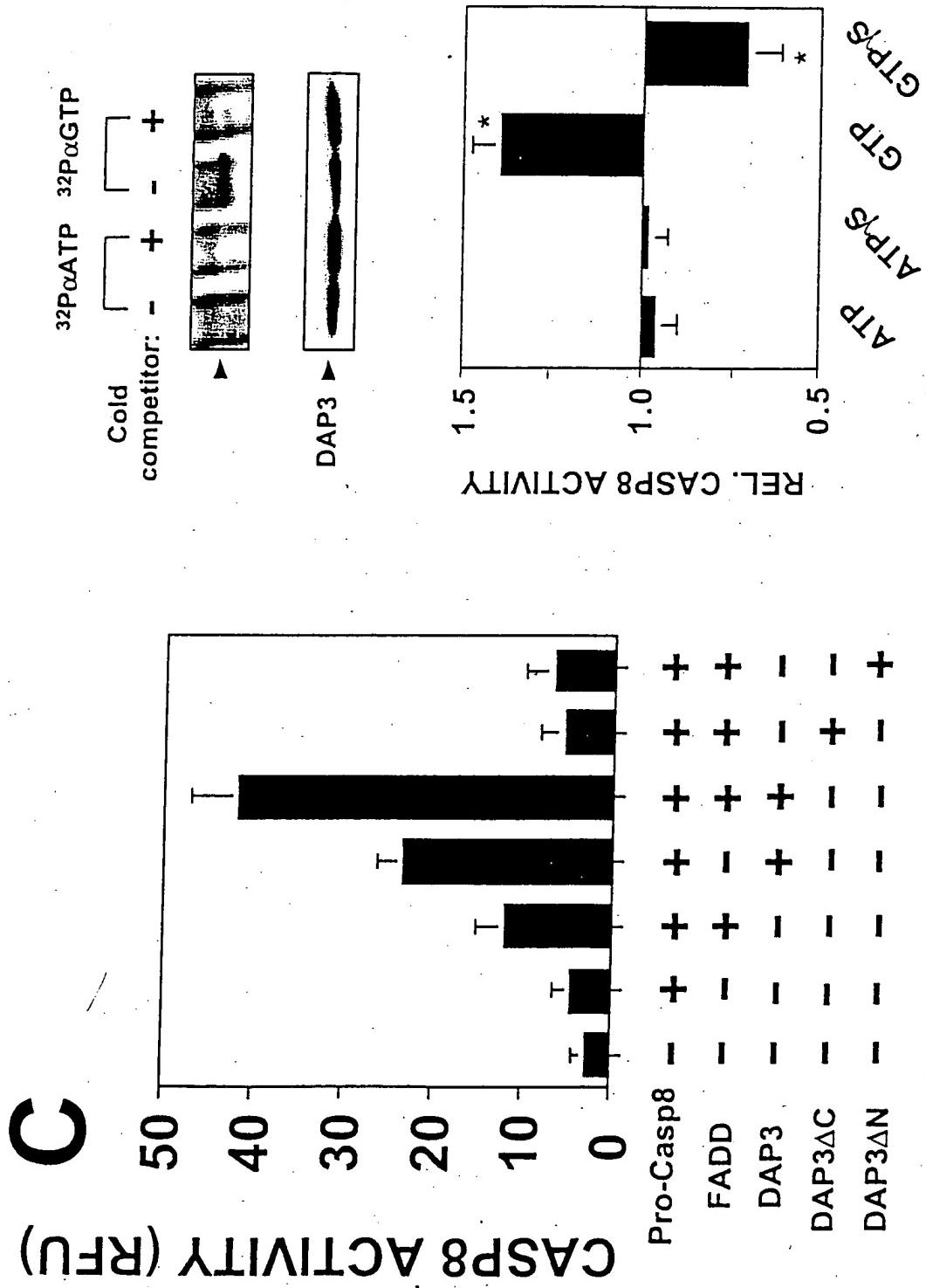


FIGURE 2C

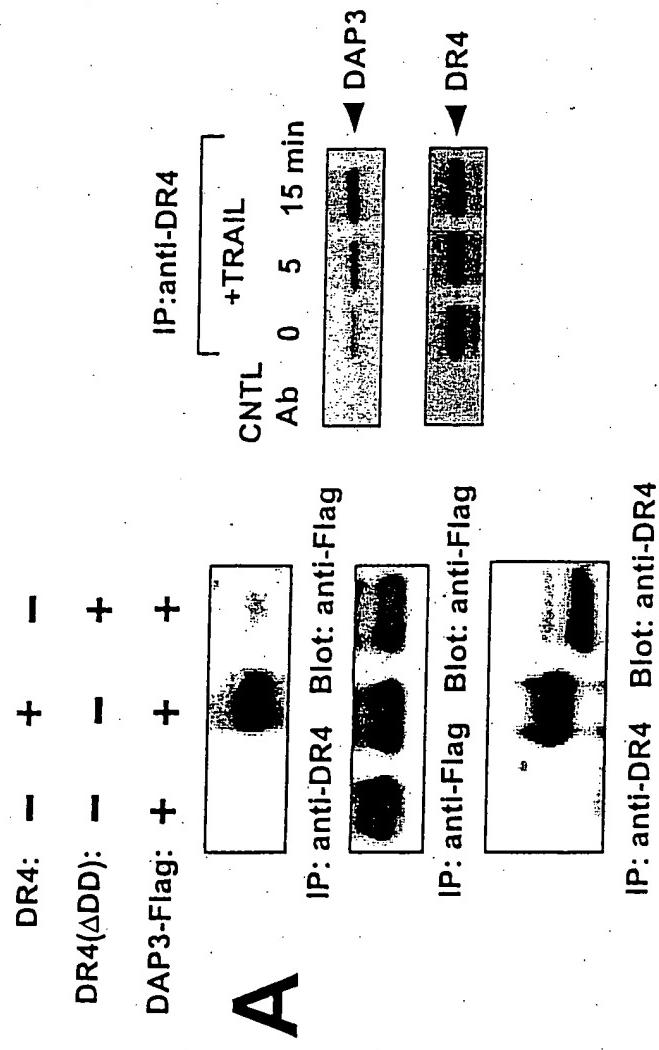


FIGURE 3A

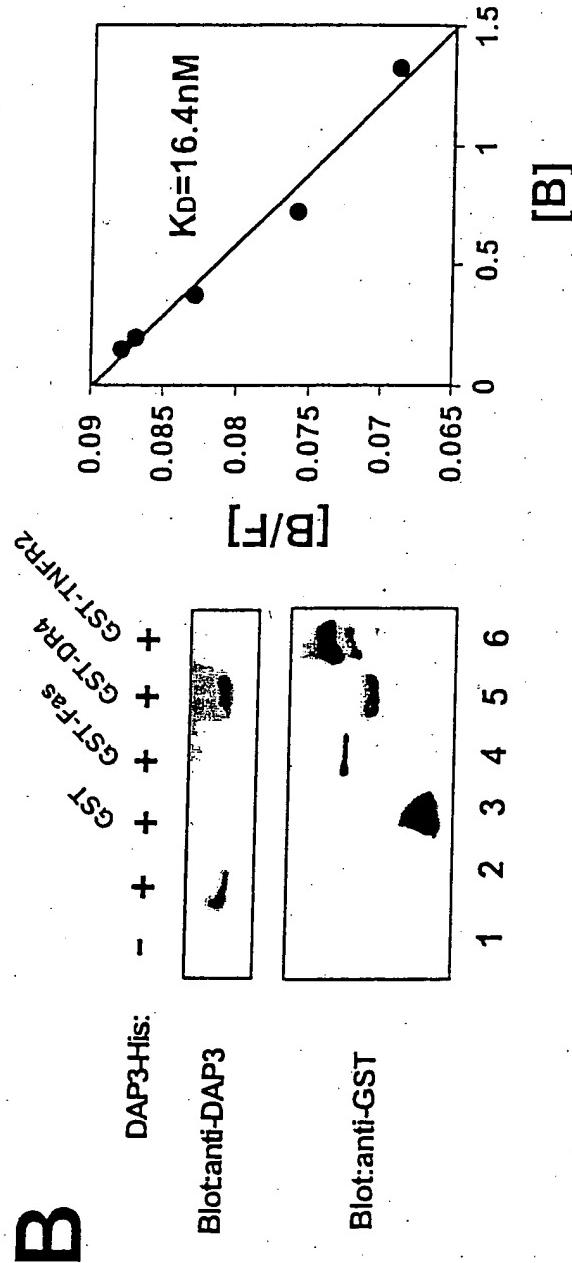


FIGURE 3B

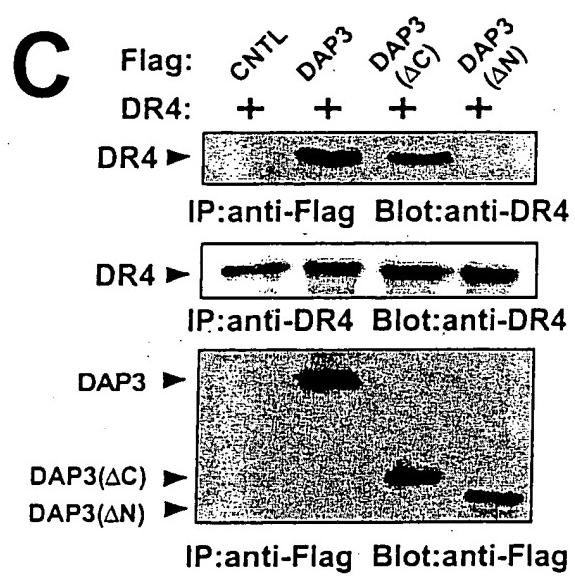


FIGURE 3C

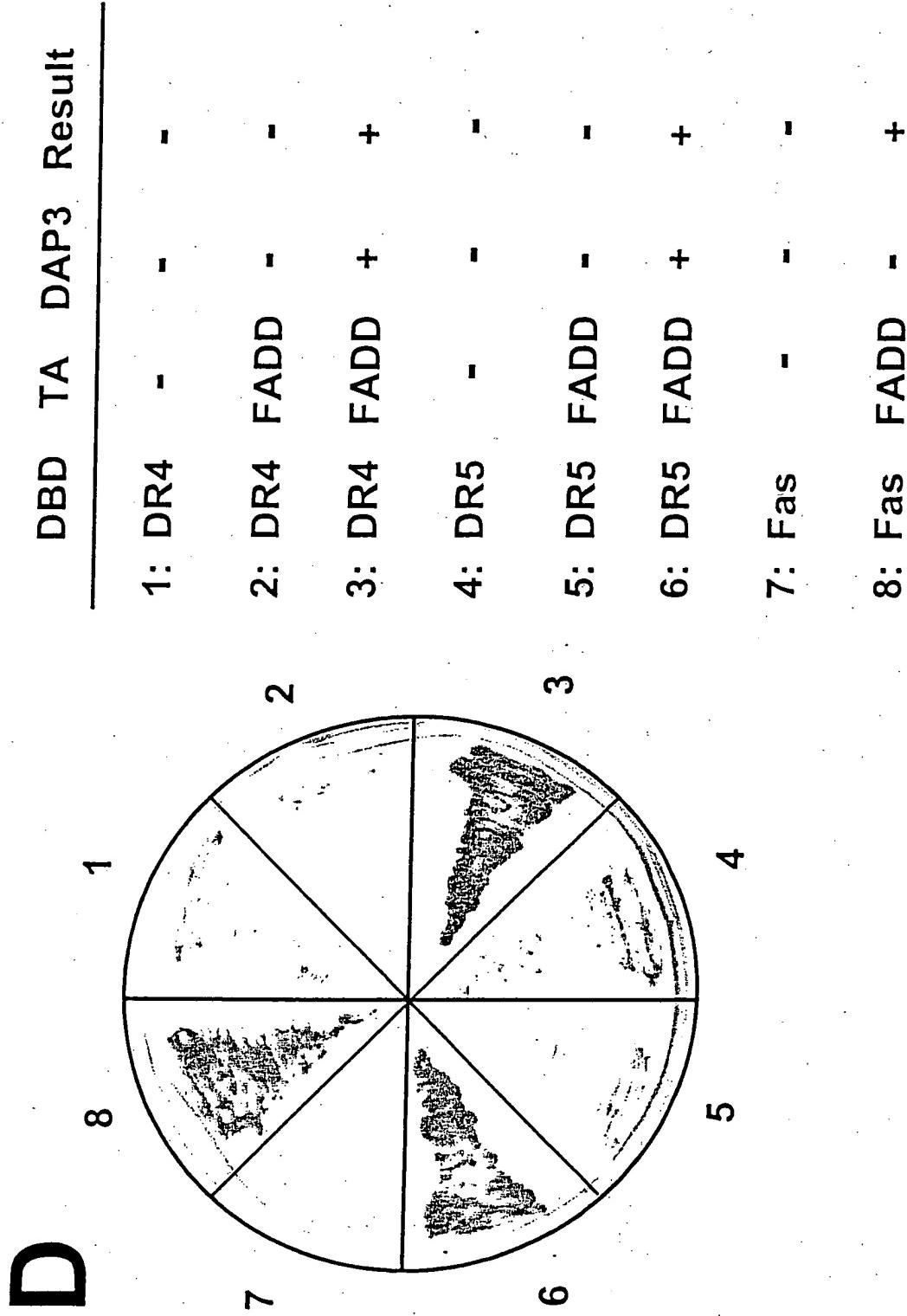


FIGURE 3D

E

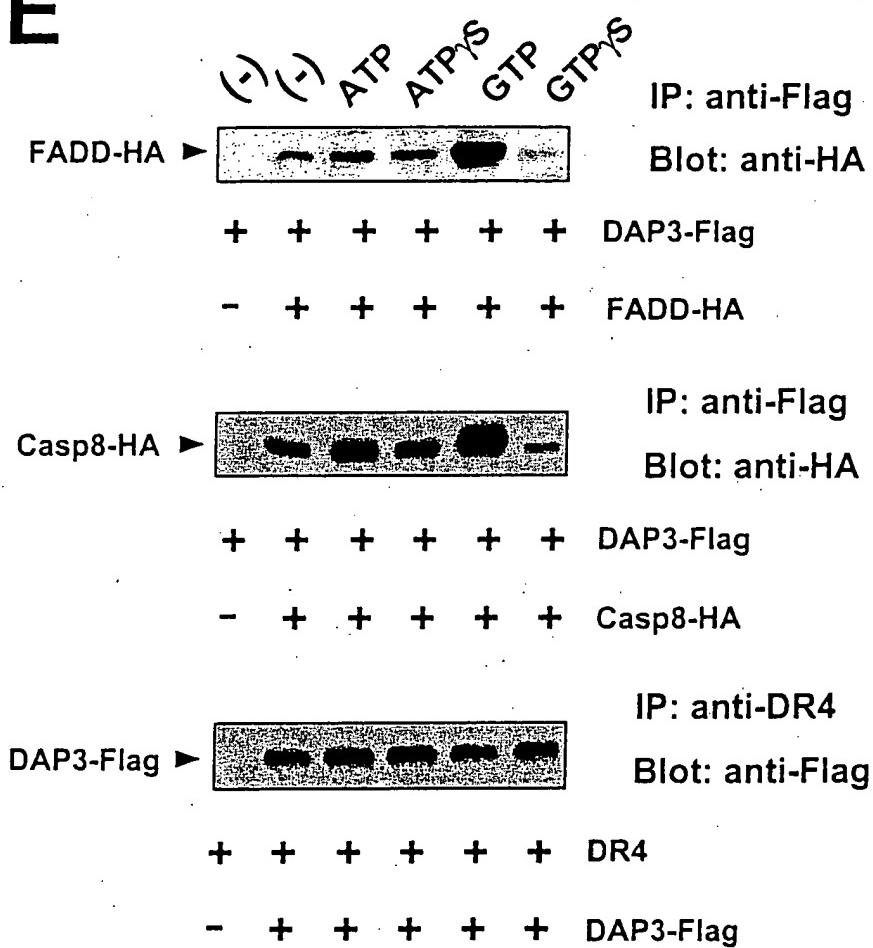


FIGURE 3E

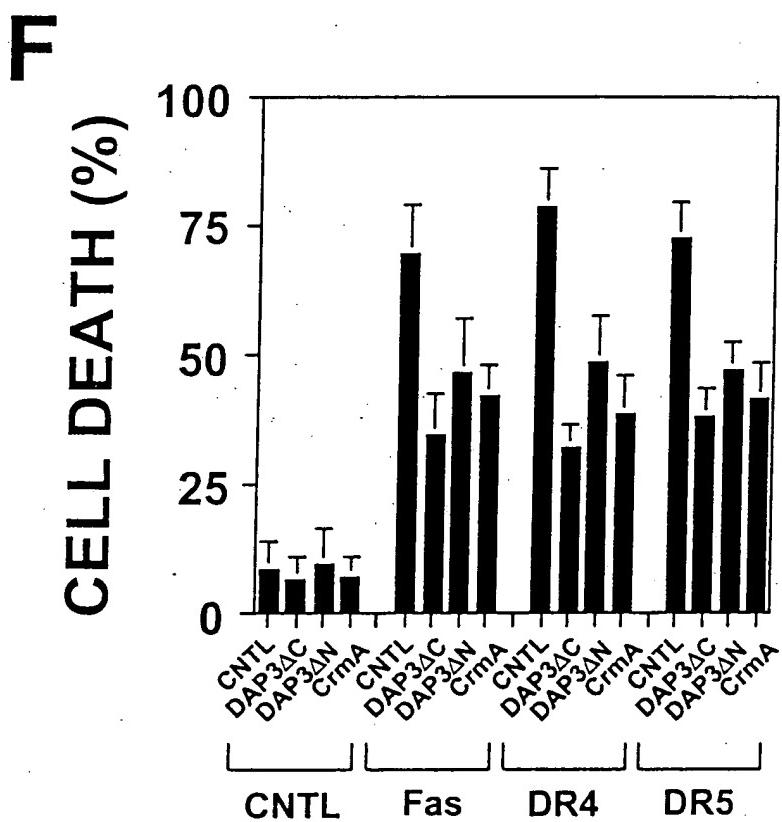


FIGURE 3F

G

DAP3
agtccctcg a ctcacgtgca aggatatgc tgaaaggaaa

antisense 1

antisense 2

antisense 3

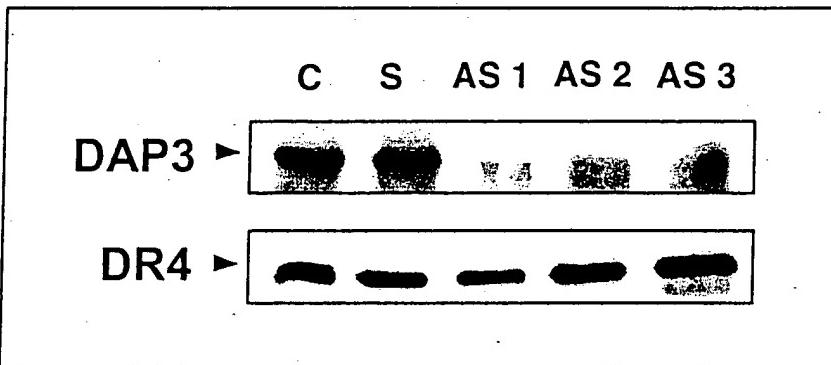
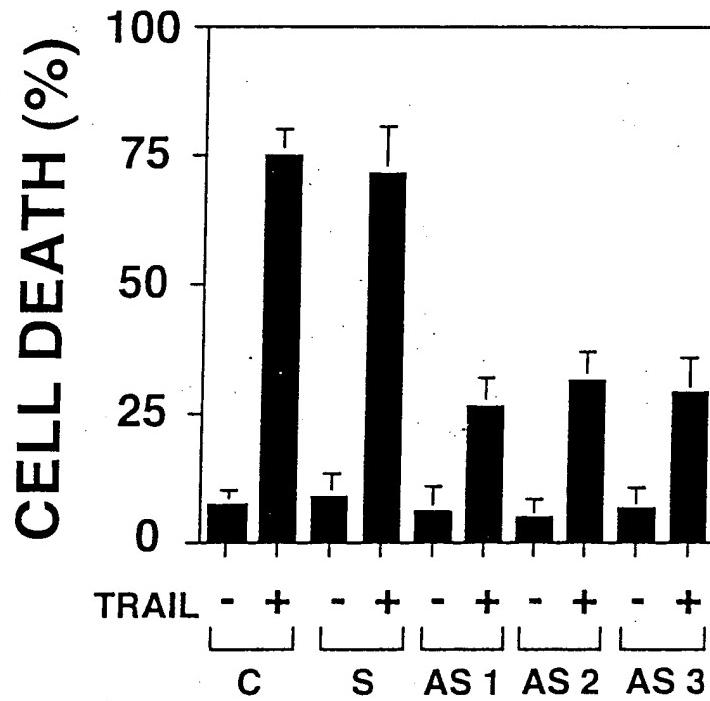


FIGURE 3G

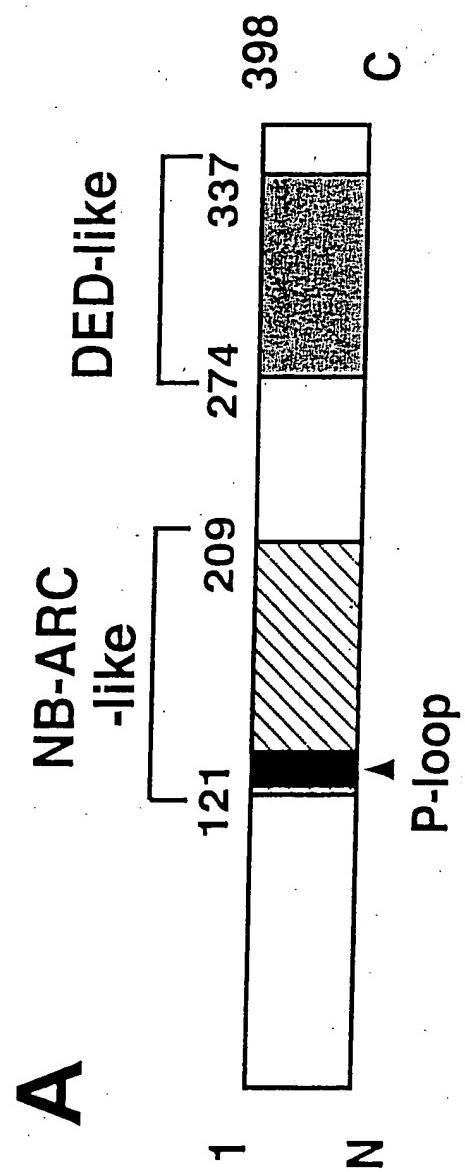


FIGURE 4A

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FIGURE 4B, C

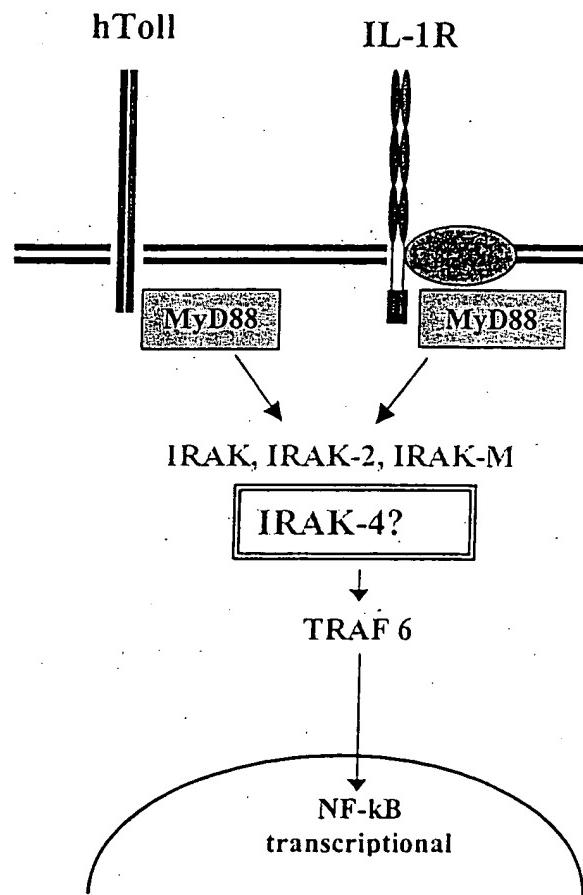


FIGURE 5

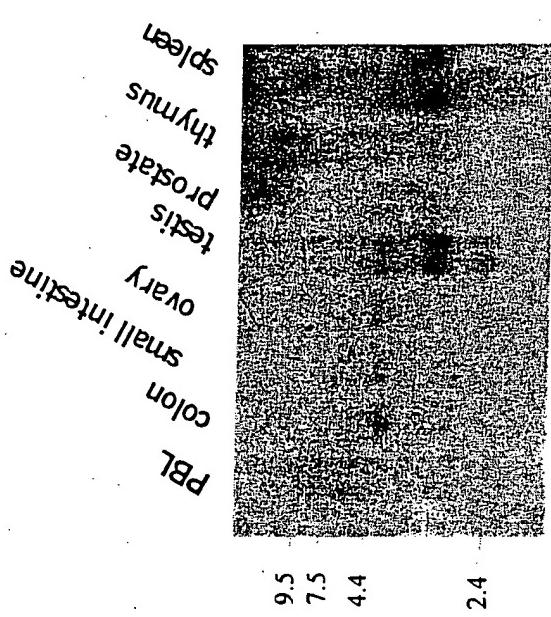


FIGURE 6

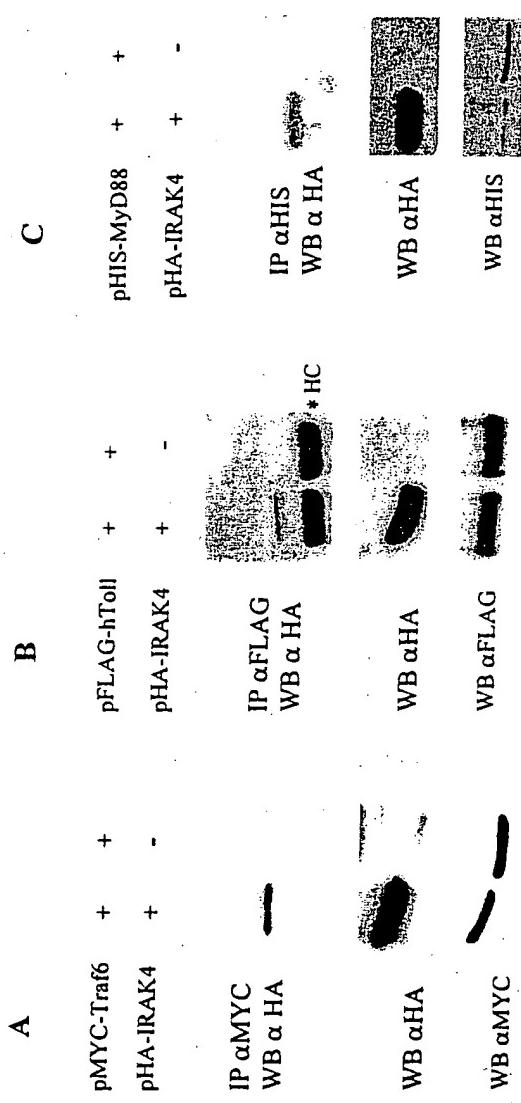


FIGURE 7

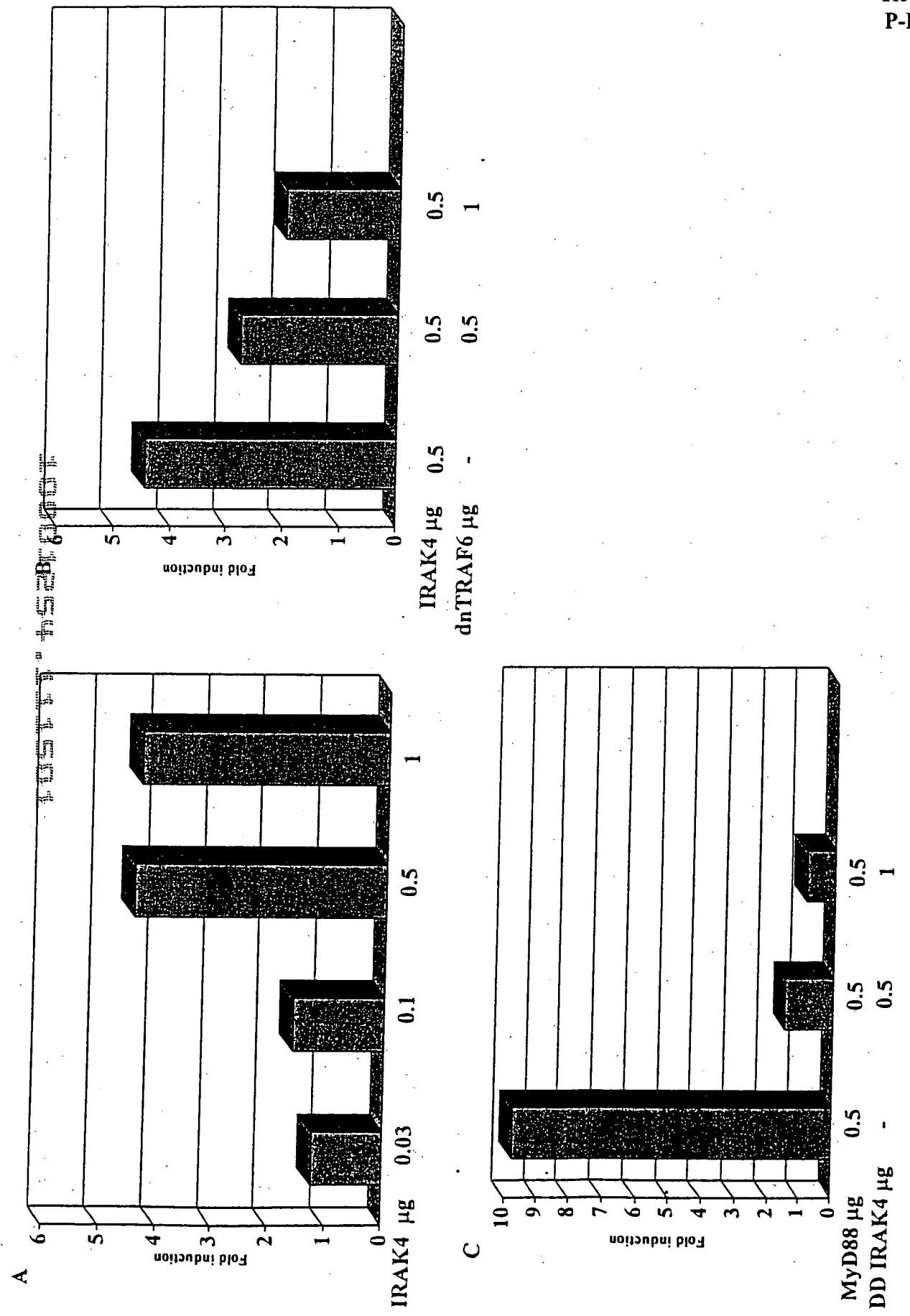


FIGURE 8

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

MEEDDECDWIGMSEH MREVAGGQTECEFEELIARSHDEA-----PG
WPEEHGEQEHGEP SHHMDDIVGTHITHDV VLSFTKIDVADP
WPEEHGEQEHGEP SHHMDDIVGTHITHDV VLSFTKIDVADP
MEETECESMETTSSEHTRTEGSQTETCGGTAEVVDDTIPGHPEDPEGSTEDPPG
MEPDWVHSVSSSISSSSTTEPACGG-----MG

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

AAGG-----IA-----ASCHEELSEEE GQCEHSNE LH
HEEG-----EN-----NGDPLAAE QGCEHSNE QG
HEEG-----EN-----NGDPLAAE QGCEHSNE QG
PEGSPQANTPCP EN SWO MOPQ EGCSIAS H P SGAELLE E GMDSAN P
EGSPQANTPCP EN SWO MOPQ EGCSIAS H P SGAELLE E GMDSAN P
EGSPQANTPCP EN SWO MOPQ EGCSIAS H P SGAELLE E GMDSAN P
EGSPQANTPCP EN SWO MOPQ EGCSIAS H P SGAELLE E GMDSAN P

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

GQSPVIAHDTPHIA PNSP
GQSPVIAHDTPHIA PNSP
GQSPVIAHDTPHIA PNSP
GQSPVIAHDTPHIA PNSP
GQSPVIAHDTPHIA PNSP

4758144_hDEDD

FIGURE 9

a tgaacaaacc cataaacacca tcaacatatg tgcgctgcct caatgttgg a ctaatttagga agctgtcaga
tttatttat cctcaagaag gatggaagaa gttagctgt a gctattaaa aaccatctgg
tcatgtataga tacaatcagt ttcacataag gagatttgaa gcattacttc aaactggaaa
aagtcccact tctgaattac tgtttgactg gggcaccaca aattgcacag ttgtgtatct
tgtggatctt ttgatccaaa atgaatttt tgctcctgct agtctttgc tccagatgc
tggcccaaa actgctaata cactaccc taaagaagct ataacagttc agaaaaaaca
gatgcctttc tgtgacaaag acaggacatt gatgacacct gtgcagaatc ttgaacaaag
ctatatgccca cctgactcct caagtccaga aaataaaaat tttagaagttt gtgatacac
tttccacagt ttgttcatat atgaatttga gaatgtcaca aataactttt atgaacgacc
catttctgtt ggtgtaata aaatgggaga gggaggattt ggagttgtat ataaaggcta
cgtaaataac acaactgtgg cagtgaagaa gcttgagca atgggtgaca ttactactga
agaactgaaa cagcagttt atcaagaaaat aaaagtaatg gcaaagtgtc aacatgaaa
cttagtagaa ctacttgggtt tctcaagtgtt tgtagatgac ctctgcttag tataatgttta
catgcctaat ggttcattgc tagacagact ctctgctt gatggactc caccacttc
ttggcacatg agatgcaaga ttgctcaggg tgcaagctt ggcataatt ttctacatga
aaatcatcat attcatagag atattaaaatg tgcaaatatc ttactggatg aagctttac
tgctaaaata tctgactttt gccttgcacg ggcttctgag aagtttgc agacagtc
gactagcaga attgtggaa caacagctt tatggcacca gaagcttgc gtggagaaat
aacacccaaa tctgatattt acagcttgg tgggttttta ctagaaataa taactggact
tccagctgtt gatgaacacc gtgaacctca gttattgcta gatattaaag aagaaattga
agatgaagaa aagacaattt aagatttat tggataaaaag atgaatgtt ctgattccac
ttcagttgaa gctatgtact ctgttgc tag tcaatgtctg catgaaaaga aaaataagag
accagacatt aagaaggttc aacagctgct gcaagagatg acagtttctt aa

FIGURE 10A

MNKPITPSTYVRCLNVGLIRKLSDFIDPQEGWKKLAVAIIKKPSG
DDRYNQFHIRRFEAALLQTGKSPTSELLFDWGTTNCTVGDLVDLLIQNEFFAPASLLP
DAVPKTANTLPSKEAITVQQKQMPFCDKDRTLMTPVQNLEQSYPDSSSPENKSLEV
SDTRFHFSFYELKNVTNNFDERPISVGGNKMGEGGFGVVYKGYVNNTTVAVKKLAAM
VDITTEELKQQFDQEIKVMAKCQHENLVELLGFSSDGDDLCLVYVYMPNGSLLDRLSC
LDGTPPPLSWHMRCKIAQGAANGINFLHENHHIHRDIKSANILLDEAFTAKISDFGLAR
ASEKFAQTVMTSRIVGTTAYMAPEALRGEITPKSDIYSFGVVLLEIITGLPAVDEHRE
PQLLDIKEEIEDEEKTIEDYIDKKMNDADSTSVEAMYSVASQCLHEKKNKRDPDIKKV
QQLQEMTAS

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FIGURE 10B

Death domains:

C. pneumoniae

DLWRQFALSLGVSEEE LANHEFSQAAQDMVATFRRLCDMPQLAVGLGALYTYEIQIPQVCVEKIR

C. muridarum

DLWKQFVFALGVSEELEAHEPSEAAKAKVATFMRWCTGDSLAAGVAALYSYESQIPCVAKEKIR

C. psittaci

DLWKNFAYALGVTEEELENHVPAAAQKKVDTFLRWCTGDSL SAGVAALYTYESQIPTVAETKIS

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FIGURE 11A

GI:7190927

ORGANISM Chlamydia muridarum
ORIGIN

1 mesrkgihev smnfldqlida iiqnkhamleh pfymkwskge ltkeqlqaya
kdyylhikaf

61 pkylsaihsr cddlearkll 1dnlmdeeng ypnhidlwkq fvfalgvsse
eleahepsea

121 akakvatfmr wctgdsllaag vaalysyesq ipcvakekir glieyfgfsn
pedyayftek

181 eeadvrhare ekaliemlsr ddsdkvleas revtqslygf ldsflepatic
chchka

GI:7468151

ORGANISM Chlamydophila pneumoniae [Chlamydia pneumoniae]
ORIGIN

1 mtswielldk qiedqhmlkh efyqrwsegk lekqqlqaya kdyylhikaf
pcylsalhar

61 cddlqirrqi lenlmdeeag npnhidlwrq falslgvsee elanhefsqa
aqdmvatfrr

121 lcdmpqlavg lgalytyeiq ipqvcvekir glkeyfgvsar gyayftvhq
eadikhasee

181 kemlqtlvgr enpdavlqgs qevldtlwnf lssfinstep csck

FIGURE 11B

6405 ttaagc tttgtgacaa
6421 tgacaacatg ttgcaggctc taaaaatgaa tccaaaaagc cgtataaaga ttgtgttaact
6481 tctcgcgaag cttctaaaac tttgtcgcta tcacatctctg acaacatctc aattaaggcc
6541 ttttcttccc tagcatgacg cacatcagct tcttcatgct ccgtgaaata agcataatct
6601 tcaggattag aaaagccaaa gtactcaatc aatccacgaa tttttcttt agctacgcaa
6661 ggaatttgac tttcataaga atacaaagcc gctactcctg ctgctaaaga atccccctgtg
6721 caccaccgca taaatgtcgc aaccttagct ttagctgctt cactgggttc atgagcttct
6781 agctcttctg aagacactcc aagagcaaac acaaattgtt tccataaaatc aatatgatta
6841 ggataaccat tctcttcatc cattaagtt tctaataata acttgccggc ttctaaatca
6901 tcacaacggc tatgaatagc agataaaat tttggaaaag ctttgatatg caaatagtaa
6961 tcttttgct atgcctgtaa ttgttcttt gtca gctctc cttttgacca cttcatgtaa
7021 aaagggtgtt ctaacatatg tttgtttga ataattgcat ctagctgatc taaaaaattc
7081 atgctcacct cttttattcc ttttcttgat tccac 7115

FIGURE 11C

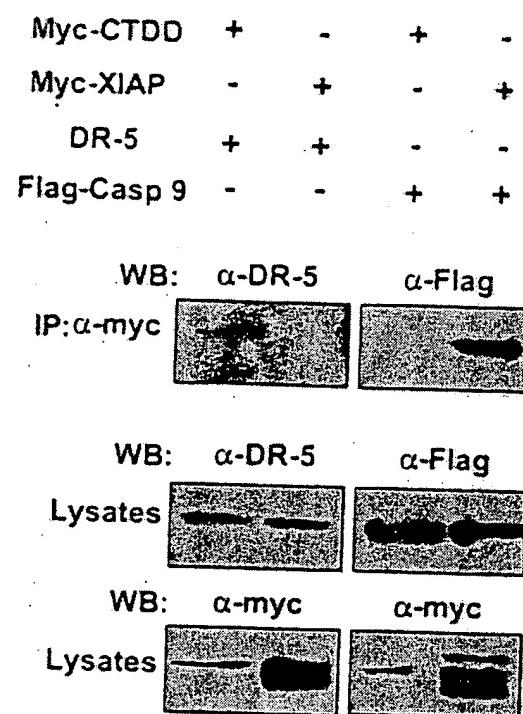


FIGURE 12

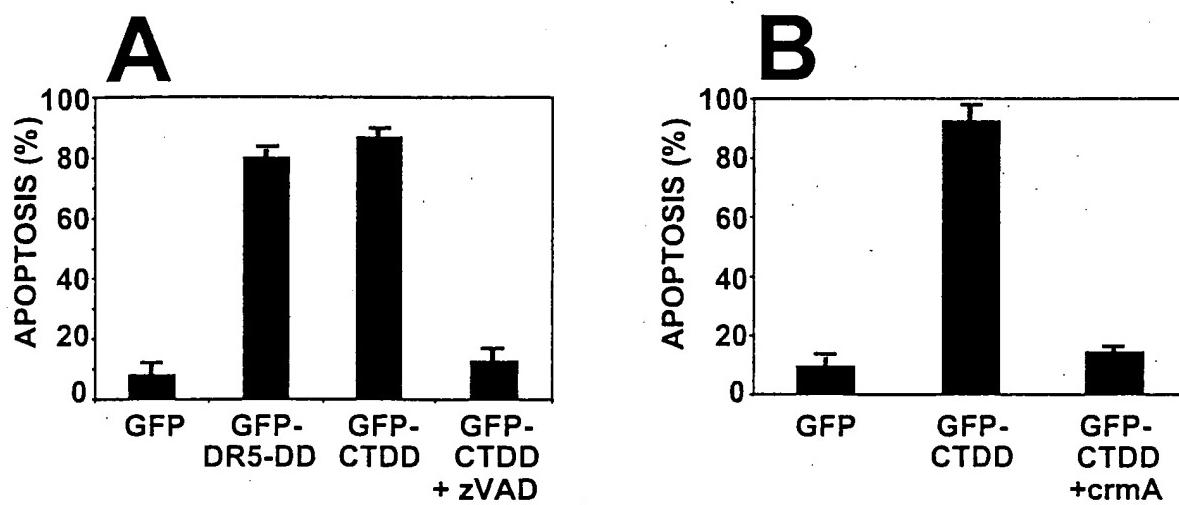


FIGURE 13

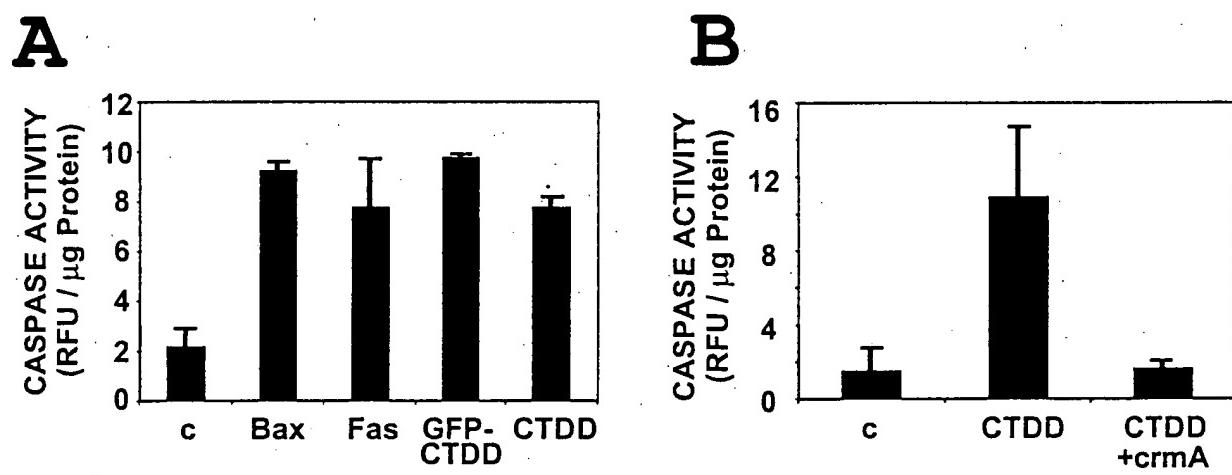


FIGURE 14

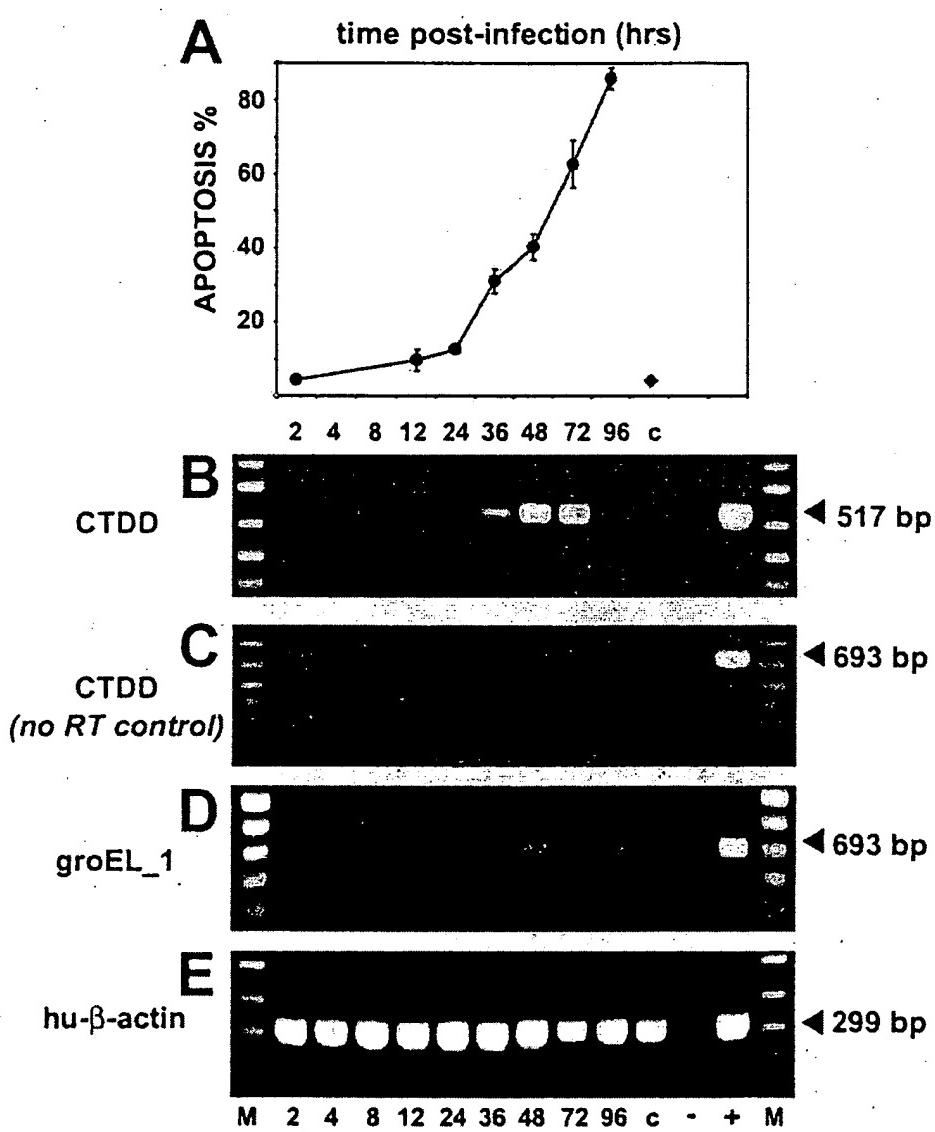


FIGURE 15